

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicant(s) : Jean-Christophe Audonnet, *et al*
U.S. Appln. No. : 09/943,443
U.S. Filing Date : 08-30-2001
Title of Invention : Feline polynucleotide vaccine formula
Confirm No. : 9956
Examiner : CHEN, STACY BROWN
Art Unit : 1648

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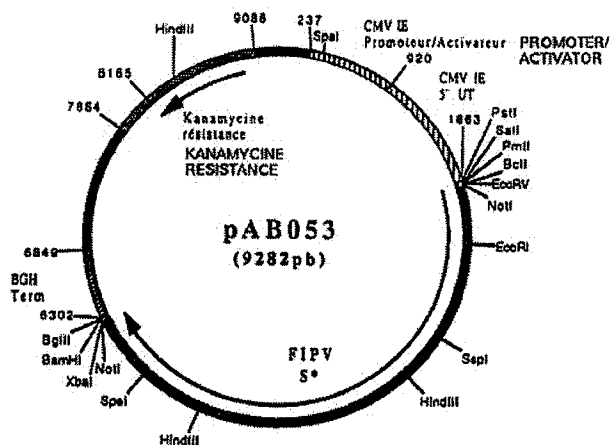
(51) Classification internationale des brevets ⁶ : C12N 15/48, 15/35, 15/50, 15/38, 15/40, 15/49, 15/47, A61K 39/295		(11) Numéro de publication internationale: WO 98/03660
A1		(43) Date de publication internationale: 29 janvier 1998 (29.01.98)
(21) Numéro de la demande internationale: PCT/FR97/01315		(81) Etats désignés: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GE, GH, HU, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ZW, brevet ARIPO (GH, KE, LS, MW, SD, SZ, UG, ZW), brevet eurasién (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), brevet européen (AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), brevet OAPI (BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG).
(22) Date de dépôt international: 15 juillet 1997 (15.07.97)		
(30) Données relatives à la priorité: 96/09337 19 juillet 1996 (19.07.96) FR		
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(54) Title: FELINE POLYNUCLEOTIDE VACCINE FORMULA

(54) Titre: FORMULE DE VACCIN POLYNUCLEOTIDIQUE FELIN

(57) Abstract

A cat vaccine formula including at least three polynucleotide vaccine valencies that each include a plasmid containing a cat pathogen valency gene capable of being expressed *in vivo* in host cells. Said valencies are selected from the group which consists of feline leukaemia virus, panleukopenia virus, infectious peritonitis virus, coryza virus, calicivirus disease virus, feline immunodeficiency virus and optionally rabies virus. Said plasmids include one or more genes per valency, and said genes are selected from the group which consists of env and gag for feline leukaemia, VP2 for panleukopenia, modified S, M and N for infectious peritonitis, gB and gD for coryza, capsid for calicivirus disease, env and gag/pro for feline immunodeficiency and G for rabies.



FELINE POLYNUCLEOTIDE VACCINE FORMULA

The present invention relates to a vaccine formula allowing the vaccination of cats against a number of pathologies. It also relates to a corresponding method of vaccination.

Associations of vaccines against certain canine viruses have already been proposed in the past.

The associations developed so far were prepared from inactivated vaccines or live vaccines and, optionally, from mixtures of such vaccines. Their development raises problems of compatibility between valencies and of stability. It is indeed necessary to ensure both the compatibility between the different vaccine valencies, whether from the point of view of the different antigens used or from the point of view of the formulations themselves, especially in the case where both inactivated vaccines and live vaccines are combined. The problem of the conservation of such combined vaccines and of their safety especially in the presence of an adjuvant also exists. These vaccines are in general quite expensive.

Patent Applications WO-A-90 11092, WO-A-93 19183, WO-A-94 21797 and WO-A-95 20660 have made use of the recently developed technique of polynucleotide vaccines. It is known that these vaccines use a plasmid capable of expressing, in the host cells, the antigen inserted into the plasmid. All routes of administration have been proposed (intraperitoneal, intravenous, intramuscular, transcutaneous, intradermal, mucosal and the like). Various vaccination means can also be used, such as DNA deposited at the surface of gold particles and projected so as to penetrate into the animals' skin (Tang et al., Nature 356, 152-154, 1992) and liquid jet injectors which make it possible to transfect at the same time the skin, the muscle, the fatty tissues and the mammary tissues (Furth et al., Analytical Biochemistry, 205, 365-368, 1992).

Polynucleotide vaccines may also use naked DNAs as well as DNAs formulated, for example, inside cationic lipids or liposomes.

The invention therefore proposes to provide a multivalent vaccine formula that makes possible to ensure vaccination against a number of feline pathogenic viruses.

Another objective of the invention is to provide such a vaccine formula combining different valencies while exhibiting all the criteria required for mutual compatibility and stability of the valencies.

Another objective of the invention is to provide such a vaccine formula which makes it possible to combine different valencies in the same vehicle.

Another objective of the invention is to provide such a vaccine which is easy and inexpensive to use.

Yet another objective of the invention is to provide a method for vaccinating cats allowing protection, including multivalent protection, with a high level of efficiency and of long duration, as well as good safety.

The subject of the present invention is therefore a vaccine formula intended for cats, comprising at least three polynucleotide vaccine valencies each comprising an integrating plasmid, so as to express it in vivo in the host cells, a gene of a feline pathogen valency, these valencies being selected from the group consisting of feline leukaemia virus (FeLV), panleukopenia virus (FPV), infectious peritonitis virus (FIPV), coryza virus (FHV), calicivirocic virus (FCV), feline immunodeficiency virus (FIV) and possibly rabies virus (rhabdovirus), the plasmids comprising, for each valency, one or more of the genes selected from the group consisting of env and gag/pol for the feline leukaemia, VP2 for the panleukopenia, modified S (or S*) and M for the infectious peritonitis, gB and gD for the coryza, capsid for the calicivirocic, env and gag/pro for the feline immunodeficiency and G for the rabies.

Valency in the present invention is understood to mean at least one antigen providing protection against the virus of the pathogen considered, wherein the valency may contain, as subvalency, one or more modified or natural genes from one or more strains of the pathogen considered.

Gene of a pathogenic agent is understood to mean not only the whole gene but also the various nucleotide sequences, including fragments which retain the ability to induce a protective response. The term gene encompasses nucleotide sequences equivalent to those precisely described in the examples, i.e. the sequences which are different but which encode the same protein. It also encompasses the nucleotide sequences of other strains of the pathogen

considered, providing a cross-protection or a protection specific for a strain or for a strain group. It also encompasses the nucleotide sequences which have been modified in order to facilitate the in vivo expression by the host animal but which encode the same protein.

Preferably, the vaccine formula according to the invention comprises the panleukopenia, coryza and calicivirosis valencies.

It will be possible to add the feline leukaemia, feline immunodeficiency and/or infectious peritonitis valencies.

As far as the coryza valency is concerned, it is preferable to use the two genes coding for gB and gD integrated in different plasmids or in a single plasmid, or to use either of these genes.

For the feline leukaemia valency, use is preferably made of the two env and gag/pol genes integrated into two different plasmids or into a single plasmid, or the env gene alone.

For the feline immunodeficiency valency, use will preferably be made of the two env and gag/pro genes in different plasmids or in a single plasmid, or only one of these genes. Still more preferably, the FeLV-A env gene and the FeLV-A and FeLV-B env genes are used.

For the infectious peritonitis valency, use is preferably made of the two M and modified S genes together in two different plasmids or in a single plasmid, or either of these genes. S will be modified in order to inactivate the major facilitating epitopes, preferably according to the teaching of Patent PCT/FR95/01128.

The vaccine formula according to the invention can be presented in a dose volume of between 0.1 and 3 ml and in particular between 0.5 and 1 ml.

The dose will be generally between 10 ng and 1 mg, preferably between 100 ng and 500 µg and still more preferably between 1 µg and 250 µg per plasmid type.

Use will preferably be made of naked plasmids simply placed in the vaccination vehicle which will be, in general, physiological saline (0.9% NaCl), ultrapure water, TE buffer and the like. All the polynucleotide vaccine forms described in the prior art can of course be used.

Each plasmid comprises a promoter capable of ensuring the expression into the host cells of the gene inserted, under its control. This will be in general a strong eukaryotic promoter

and in particular a cytomegalovirus early CMV-IE promoter of human or murine origin, or optionally of another origin such as rats, pigs and guinea pigs.

More generally, the promoter may be either of viral origin or of cellular origin. As viral promoter, there may be mentioned the SV40 virus early or late promoter or the Rous sarcoma virus LTR promoter. It may also be a promoter from the virus from which the gene is derived, for example the gene's own promoter.

As cellular promoter, there may be mentioned the promoter of a cytoskeleton gene, such as for example the desmin promoter (Bolmont et al., *Journal of Submicroscopic Cytology and Pathology*, 1990, 22, 117-122; and Zhenlin et al., *Gene*, 1989, 78, 243-254), or alternatively the actin promoter.

When several genes are present in the same plasmid, these may be presented in the same transcription unit or in two different units.

The combination of the different vaccine valencies according to the invention may preferably be achieved by mixing the polynucleotide plasmids expressing the antigen(s) of each valency, but it is also possible to have the same plasmid express antigens of several valencies.

The subject of the invention is also monovalent vaccine formulae comprising one or more plasmids encoding one or more genes from one of the viruses cited above, the genes being those described above. Besides their monovalent character, these formulae may possess the characteristics stated above regarding the choice of the genes, their combinations, the composition of the plasmids, the dose volumes, the doses and the like.

The monovalent vaccine formulae may also be used (i) for the preparation of a polyvalent vaccine formula as described above, (ii) individually against the actual pathology, (iii) combined with a vaccine of another type (live or inactivated whole, recombinant, subunit) against another pathology, or (iv) as booster for a vaccine as described below.

The subject of the present invention is in fact also the use of one or more plasmids according to the invention for the manufacture of a vaccine intended to vaccinate cats primo-vaccinated by means of a first conventional vaccine (monovalent or multivalent) as known in the prior art, in particular, selected from the group consisting of a live whole vaccine, an inactivated whole vaccine, a subunit vaccine, a recombinant vaccine, this first vaccine having (i.e.,

containing or capable of expressing) the antigen(s) encoded by the plasmid(s) or antigen(s) providing cross-protection.

Remarkably, the polynucleotide vaccine has a potent booster effect which results in an amplification of the immune response and the acquisition of a long-lasting immunity.

In general, the primo-vaccination vaccines can be selected from commercial vaccines available from various veterinary vaccine producers.

The subject of the invention is also a vaccination kit assembling together a primo-vaccination vaccine as described above and a vaccine formula according to the invention for the booster. It also relates to a vaccine formula according to the invention accompanied by a notice indicating the use of this formula as a booster for a primo-vaccination as described above.

The subject of the present invention is also a method for vaccinating cats, comprising the administration of an effective vaccine formula as described above. This vaccination method comprises the administration of one or more doses of the vaccine formula, wherein these doses can be administered successively over a short period of time and/or successively at widely spaced intervals.

The vaccine formulae according to the invention can be administered in the context of this method of vaccination, by different routes of administration proposed in the prior art for polynucleotide vaccination and by means of known techniques of administration.

The subject of the invention is also a method of vaccination consisting in performing a primo-vaccination as described above and a booster with a vaccine formula according to the invention.

In a preferred embodiment of the process according to the invention, in a first instance, an animal is administered an effective dose of the vaccine of the conventional, especially inactivated, live, attenuated or recombinant, type, or alternatively a subunit vaccine, so as to provide a primo-vaccination, and, after a period preferably of 2 to 6 weeks, is administered the polyvalent or monovalent vaccine according to the invention.

The efficiency of presentation of the antigens to the immune system varies according to the tissues. In particular, the mucous membranes of the respiratory track serve as barriers to the entry of pathogens and are associated with lymphoid tissues which support local immunity.

Administration of a vaccine by contact with the mucous membranes, in particular the buccal mucous membrane, the pharyngeal mucous membrane and the mucous membrane of the bronchial region, is of great interest for the vaccination against respiratory and digestive pathologies.

Consequently, administration using the mucosal routes is a preferred mode of administration for the invention, using in particular nebulization or spray or drinking water. It will be possible to apply the vaccine formulae and the vaccination methods according to the invention in this context.

The invention also relates to a method of preparing the vaccine formulae, namely the preparation of the valencies and mixtures thereof, as evident from this description.

The invention will now be described in greater detail with the aid of the embodiments of the invention taken with reference to the accompanying drawings.

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Figure No. 9:	Plasmid pAB052
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Sequence listing SEQ ID No.

SEQ ID No. 1:	Oligonucleotide PB247
SEQ ID No. 2	Oligonucleotide PB249
SEQ ID No. 3:	Oligonucleotide PB281
SEQ ID No. 4:	Oligonucleotide PB282
SEQ ID No. 5:	Sequence of the FELV-B virus env gene
SEQ ID No. 6:	Oligonucleotide PB283
SEQ ID No. 7:	Oligonucleotide PB284
SEQ ID No. 8:	Sequence of the FeLV-A virus gag/pol gene (Glasgow-1 strain)
SEQ ID No. 9:	Oligonucleotide AB021
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SEQ ID No. 11:	Oligonucleotide AB103
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SEQ ID No. 19:	Oligonucleotide AB061
SEQ ID No. 20:	Oligonucleotide AB064
SEQ ID No. 21:	Oligonucleotide AB065
SEQ ID No. 22:	Oligonucleotide AB066
SEQ ID No. 23:	Oligonucleotide AB025
SEQ ID No. 24:	Oligonucleotide AB026
SEQ ID No. 25:	Oligonucleotide AB067
SEQ ID No. 26:	Oligonucleotide AB070
SEQ ID No. 27:	Oligonucleotide AB154
SEQ ID No. 28:	Oligonucleotide AB155
SEQ ID No. 29:	Oligonucleotide AB011
SEQ ID No. 30:	Oligonucleotide AB012

EXAMPLES**Example 1: Culture of the Viruses**

The viruses are cultured on the appropriate cellular system until a cytopathic effect is obtained. The cellular systems to be used for each virus are well known to persons skilled in the art. Briefly, the cells sensitive to the virus used, cultured in Eagle's minimum essential medium (MEM medium) or another appropriate medium, are inoculated with the viral strain studied using a multiplicity of infection of 1. The infected cells are then incubated at 37°C. for a time necessary for the appearance of a complete cytopathic effect (on average 36 hours).

Example 2: Extraction of the Viral Genomic DNAs

After culture, the supernatant and the lysed cells are harvested and the entire viral suspension is centrifuged at 1000 g for 10 minutes at +4°C so as to remove the cellular debris. The viral particles are then harvested by ultracentrifugation at 400,000 g for 1 hour at +4°C. The pellet is taken up in a minimum volume of buffer (10 mM Tris, 1 mM EDTA). This concentrated viral suspension is treated with proteinase K (100 µg/ml final) in the presence of sodium dodecyl sulphate (SDS) (0.5% final) for 2 hours at 37°C. The viral DNA is then extracted with a mixture of phenol/chloroform and then precipitated with 2 volumes of absolute ethanol. After leaving the sample overnight at -20°C, the DNA is centrifuged at 10,000 g for 15 minutes at +4°C. The DNA pellet is dried and then taken up in a minimum volume of sterile ultrapure water. It can then be digested with restriction enzymes.

Example 3: Isolation of the Viral Genomic RNAs

The RNA viruses were purified according to techniques well known to one skilled in the art. The genomic viral RNA of each virus was then isolated using the "guanidium thiocyanate/phenol-chloroform" extraction technique described by P. Chomczynski and N. Sacchi (Anal. Biochem., 1987, 162, 156-159).

Example 4: Molecular Biology Techniques

All the constructions of plasmids were carried out using standard molecular biology techniques described by J. Sambrook et al. (Molecular Cloning: A Laboratory Manual, 2nd Edition, Cold

Spring Harbor Laboratory, Cold Spring Harbor, N.Y., 1989). All the restriction fragments used for the present invention were isolated using the "GeneClean" kit (BIO 101 Inc. La Jolla, Calif.).

Example 5: RT-PCR Technique

Specific oligonucleotides (comprising restriction sites at their 5' ends to facilitate the cloning of the amplified fragments) were synthesized such that they completely cover the coding regions of the genes to be amplified (see specific examples). The reverse transcription (RT) reaction and polymerase chain reaction (PCR) were carried out according to standard techniques (Sambrook J. et al., 1989). Each RT-PCR reaction was performed with a pair of specific amplimers and using, as template, the extracted viral genomic RNA. The complementary DNA amplified was extracted with phenol/chloroform/isoamyl alcohol (25:24:1) before being digested with restriction enzymes.

Example 6: Plasmid pVR1012

The plasmid pVR1012 (Figure No. 1) was obtained from Vical Inc., San Diego, Calif., USA. Its construction has been described in J. Hartikka et al. (Human Gene Therapy, 1996, 7, 1205-1217).

Example 7: Construction of the Plasmid pPB179 (FeLV-A Virus Env Gene)

An RT-PCR reaction according to the technique described in Example 5 was carried out with feline leukaemia virus (FeLV-A) (Glasgow-1 strain) genomic RNA (M. Stewart et al. J. Virol. 1986. 58. 825-834), prepared according to the technique described in Example 3, and with the following oligonucleotides:

PB247 (29 mer) (SEQ ID No. 1)

5'TTTGTCGACCATGGAAAGTCCAACGCACC3'

PB249 (28 mer) (SEQ ID No. 2)

5'TTTGGATCCTCATGGTCGGTCCGGATCG3'

so as to amplify a 1947 bp fragment containing the gene encoding the Env glycoprotein from the FeLV-A virus (Glasgow-1 strain) in the form of a SalI-BamHI fragment. After purification, the RT-PCR product was digested with SalI and BamHI in order to give a 1935 bp SalI-BamHI fragment.

This fragment was ligated with the vector pVR1012 (Example 6), previously digested with Sall and BamHI, to give the plasmid pPB179 (6804 bp) (Figure No. 2).

Example 8: Construction of the Plasmid pPB180 (FeLV-S Virus Env Gene)

An RT-PCR reaction according to the technique described in Example 5 was carried out with feline leukaemia virus (FeLV-B subtype) genomic RNA, prepared according to the technique described in Example 3, and with the following oligonucleotides:

PB281 (29 mer) (SEQ ID No. 3)

5'TTTGTCGACATGGAAGGTCCAACGCACCC3'

PB282 (32 mer) (SEQ ID No. 4)

5'TTGGATCCTCATGGTCGGTCCGGATCATATTG3'

so as to amplify a 2005 bp fragment containing the gene encoding the Env glycoprotein from the FeLV-B virus (Figure No. 3 and SEQ ID No. 5) in the form of a Sall-BamHI fragment. After purification, the RT-PCR product was digested with Sall and BamHI in order to give a 1995 bp Sall-BamHI fragment.

This fragment was ligated with the vector pVR1012 (Example 6), previously digested with Sall and BamHI, to give the plasmid pPB180 (6863 bp) (Figure No. 4).

Example 9: Construction of the Plasmid pPB181 (FeLV gag/pol gene)

An RT-PCR reaction according to the technique described in Example 5 was carried out with the feline leukaemia virus (FeLV-A subtype) (Glasgow-1 strain) genomic RNA, prepared according to the technique described in Example 3, and with the following oligonucleotides:

PB283 (33 mer) (SEQ ID No. 6)

5'TTGTGCGACATGTCTGGAGCCTCTAGTGGGACAG3'

PB284 (42 mer) (SEQ ID No. 7)

5'TTGGATCCTTATTTAATTACTGCAGTTCCAAGGAACTCTC3'

so as to amplify a 3049 bp fragment containing the sequence encoding the Gag protein and the 5' part of the sequence encoding the Pol protein from the FeLV-A virus (Glasgow-1 strain) (Figure

No. 5 and SEQ ID No. 8) in the form of a Sall-BamHI fragment. After purification, the RT-PCR product was digested with Sall and BamHI to give a 3039 bp Sall-BamHI fragment.

This fragment was ligated with the vector pVR1012 (Example 6), previously digested with Sall and BamHI, to give the plasmid pPB181 (7908 bp) (Figure No. 6).

Example 10: Construction of the Plasmid pAB009 (FPV VP2 gene)

A PCR reaction was carried out with the feline panleukopaenia virus (193 strain) genomic DNA (J. Martyn et al., J. Gen. Virol. 1990, 71. 2747-2753), prepared according to the technique of Example 2, and with the following oligonucleotides:

AB021 (34 mer) (SEQ ID No. 9)

5'TGCTCTAGAGCAATGAGTGATGGAAGCAGTTCAAC3'

AB024 (33 mer) (SEQ ID No. 10)

5'CGCGGATCCATTAATATAATTTTCTAGGTGCTA3'

so as to amplify a 1776 bp fragment containing the gene encoding the FPV VP2 capsid protein. After purification, the PCR product was digested with XbaI and BamHI in order to give a 1764 bp XbaI-BamHI fragment.

This fragment was ligated with the vector pVR1012 (Example 6), previously digested with XbaI and BamHI, to give the plasmid pAB009 (6664 bp) (Figure No. 7).

Example 11: Construction of the Plasmid pAB053 (FIPV S* gene)

An RT-PCR reaction according to the technique described in Example 5 was carried out with the feline infectious peritonitis (FIP) virus (79-1146 strain) genomic RNA (R. de Groot et al., J. Gen. Virol. 1987. 68. 2639-2646), prepared according to the technique described in Example 3, and with the following oligonucleotides:

AB103 (38 mer) (SEQ ID No. 11)

5'ATAAGAATGCGGCCGCATGATTGTGCTCGTAACTTGCC3'

AB112 (25 mer) (SEQ ID No. 12)

5'CGTACATGTGGAATCCACTGGTTG3'

so as to amplify the sequence of the 5' part of the gene encoding the virus S glycoprotein in the form of an NotI-EcoRI fragment. After purification, the 492 bp RT-PCR product was digested with NotI and EcoRI in order to liberate a 467 bp NotI-EcoRI fragment (fragment A).

The plasmid pJCA089 (Patent Application PCT/FR95/01128) was digested with EcoRI and SpeI in order to liberate a 3378 bp fragment containing the central part of the gene encoding the FIP virus modified S glycoprotein (fragment B).

An RT-PCR reaction according to the technique described in Example 5 was carried out with the FIP virus (79-1146 strain) genomic RNA, prepared according to the technique described in Example 3, and with the following oligonucleotides:

AB113 (25 mer) (SEQ ID No. 13)

5'AGAGTTGCAACTAGTTCTGATTTTG3'

AB104 (37 mer) (SEQ ID No. 14)

5'ATAAGAATGCGGCCGCTTAGTGGACATGCACTTTTTC3'

so as to amplify the sequence of the 3' part of the gene encoding the FIP virus S glycoprotein in the form of an SpeI-NotI fragment. After purification, the 543 bp RT-PCR product was digested with SpeI and NotI in order to liberate a 519 bp SpeI-NotI fragment (fragment C).

Fragments A, B and C were then ligated together into the vector pVR1012 (Example 6), previously digested with NotI, to give the plasmid pAB053 (9282 bp), which contains the modified S gene in the correct orientation relative to the promoter (Figure No. 8).

Example 12: Construction of the Plasmid pAB052 (FIPV M gene)

An RT-PCR reaction according to the technique described in Example 5 was carried out with the feline infectious peritonitis (FIP) virus (79-1146 strain) genomic RNA (H. Vennema et al., Virology. 1991, 181. 327-335), prepared according to the technique described in Example 3, and with the following oligonucleotides:

AB101 (37 mer) (SEQ ID No. 15)

5'ACGCGTCGACCCACCATGAAGTACATTTTGCTAATAC3'

AB102 (36 mer) (SEQ ID No. 16)

5'CGCGGATCCTTACACCATATGTAATAATTTTTCATG3'

so as to precisely isolate the gene encoding the FIP virus M glycoprotein in the form of a Sall-BamHI fragment. After purification, the 812 bp RT-PCR product was digested with Sall and BamHI in order to liberate a 799 bp Sall-BamHI fragment. This fragment was then ligated into the vector pVR1012 (Example 6), previously digested with Sall and BamHI, to give the plasmid pAB052 (5668 bp) (Figure No. 9).

Example 13: Construction of the Plasmid pAB056 (FIPV N gene)

An RT-PCR reaction according to the technique described in Example 5 was carried out with the feline infectious peritonitis (FIP) virus (79-1146 strain) genomic RNA (H. Vennema et al., Virology. 1991, 181. 327-335), prepared according to the technique described in Example 3, and with the following oligonucleotides:

AB106 (35 mer) SEQ ID No. 17)

5'ACGCGTCGACGCCCTGGCCACACAGGGACAACGCG3'

AB107 (36 mer) (SEQ ID No. 18)

5'CGCGGATCCTTAGTCGTAACCTCATCAATCATCTC3'

so as to precisely isolate the gene encoding the FIP virus N protein in the form of a Sall-BamHI fragment. After purification, the 1156 bp RT-PCR product was digested with Sall and BamHI in order to liberate a 1143 bp Sall-BamHI fragment. This fragment was then ligated into the vector pVR1012 (Example 6), previously digested with Sall and BamHI, to give the plasmid pAB056 (6011 bp) (Figure No. 10).

Example 14: Construction of the Plasma pAB028 (FEV gB gene)

A PCR reaction was carried out with the feline herpesvirus (FHV-1) (C27 strain) genomic DNA (S. Spatz et al. Virology. 1993. 197. 125-36) prepared according to the technique of Example 2, and with the following oligonucleotides:

AB061 (36 mer) (SEQ ID No. 19)

5'AAAACTGCAGAATCATGTCCACTCGTGGCGATCTTG3'

AB064 (40 mer) (SEQ ID No. 20)

5'ATAAGAATGCGGCCCTTAGACAAGATTTGTTTCAGTATC3'

so as to amplify a 2856 bp fragment containing the gene encoding the FHV-1 virus gB glycoprotein in the form of a PstI-NotI fragment. After purification, the PCR product was digested with PstI and NotI to give a 2823 bp PstI-NotI fragment.

This fragment was ligated with the vector pVR1012 (Example 6), previously digested with PstI and NotI, to give the plasmid pAB028 (7720 bp) (Figure No. 11).

Example 15: Construction of the Plasmid pAB029 (FHV gD gene)

A PCR reaction was carried out with the feline herpesvirus (FHV-1) (C-27 strain) genomic DNA (S. Spatz et al. J. Gen. Virol. 1994. 75. 1235-1244), prepared according to the technique described in Example 2 and with the following oligonucleotides:

AB065 (36 mer) (SEQ ID No. 21)

5'AAAACTGCAGCCATGATGACACGTCTACATTTTGTG3'

AB066 (33 mer) (SEQ ID No. 22)

5'GGAAGATCTTTAAGGATGGTGAGTTGTATGTAT3'

so as to amplify the gene encoding the FHV-1 virus gD glycoprotein in the form of a PstI-BglII fragment. After purification, the 1147 bp PCR product was digested with PstI and BglII in order to isolate a 1129 bp PstI-BglII fragment. This fragment was ligated with the vector pVR1012 (Example 6), previously digested with PstI and BglII, to give the plasmid pAB029 (5982 bp) (Figure No. 12).

Example 16: Construction of the Plasmid pAB010 (FCV C gene)

An RT-PCR reaction according to the technique described in Example 5 was carried out with the feline calicivirus (FCV) (F9 strain) genomic RNA (M. Carter et al. Virology. 1992. 190. 443-448), prepared according to the technique described in Example 3, and with the following oligonucleotides:

AB025 (33 mer) (SEQ ID No. 23)

5'ACGCGTCGACGCATCTGCTCAACCTGCGCTAAC3'

AB026 (31 mer) (SEQ ID No. 24)

5'CGCGGATCCTCATAACTTAGTCATGGGACTC3'

so as to isolate the gene encoding the FCV virus capsid protein in the form of a SalI-BamHI fragment. After purification, the 2042 bp RT-PCR product was digested with SalI and BamHI in order to isolate a 2029 bp SalI-BamHI fragment. This fragment was ligated with the vector PVR1012 (Example 6), previously digested with SalI and BamHI, to give the plasmid pAB010 (6892 bp) (Figure No. 13).

Example 17: Construction of the Plasmid pAB030 (FIV env gene)

An RT-PCR reaction according to the technique of Example 5 was carried out with the feline immunodeficiency virus (FIV) (Petaluma strain) genomic RNA (R. Olmsted, et al. Proc. Natl. Acad. Sci. USA. 1989. 86. 8083-8096), prepared according to the technique of Example 3, and with the following oligonucleotides:

AB067 (36 mer) (SEQ ID No. 25)

5'AAAACCTGCAGAAGGAATGGCAGAAGGATTTGCAGCC3'

AB070 (36 mer) (SEQ ID No. 26)

5'CGCGGATCCTCATTCTCCTCTTTTCAGACATGCC3'

so as to amplify a 2592 bp fragment containing the gene encoding the Env glycoprotein from the FIV virus (Petaluma strain) in the form of a PstI-BamHI fragment. After purification, the RT-PCR product was digested with PstI and BamHI to give a 2575 bp PstI-BamHI fragment.

This fragment was ligated with the vector pVR1012 (Example 6), previously digested with PstI and BamHI, to give the plasmid pAB030 (7436 bp) (Figure No. 14).

Example 18: Construction of the Plasmid pAB083 (FIV gag/pro gene)

An RT-PCR reaction according to the technique described in Example 5 was carried out with the feline immunodeficiency virus (FIV) (Petaluma strain) genomic RNA (R. Olmsted et al. Proc. Natl. Acad. Sci. USA. 1989. 86. 8088-8096), prepared according to the technique described in Example 3, and with the following oligonucleotides:

AB154 (32 mer) (SEQ ID No. 27)

5'ACGCGTCGACATGGGGAATGGACAGGGGCGAG3'

AB155 (33 mer) (SEQ ID No. 28)

5'TGAAGATCTTCACTCATCCCCTTCAGGAAGAGC3'

so as to amplify a 4635 bp fragment containing the gene encoding the Gag and Pro proteins from the FIV virus (Petaluma strain) in the form of a SalI-BglII fragment. After purification, the RT-PCR product was digested with SalI and BglII to give a 4622 bp SalI-BglII fragment.

This fragment was ligated with the vector pVR1012 (Example 6), previously digested with SalI and BglII, to give the plasmid pAB083 (7436 bp) (Figure No. 15).

Example 19: Construction of the Plasmid pAB041 (rabies virus G gene)

An RT-PCR reaction according to the technique described in Example 5 was carried out with the rabies virus (ERA strain) genomic RNA (A. Anilionis et al. Nature. 1981. 294. 275-278), prepared according to the technique described in Example 3, and with the following oligonucleotides:

AB011 (33 mer) (SEQ ID No. 29)

5'AAAACCTGCAGAGATGGTTCCTCAGGCTCTCCTG3'

AB012 (34 mer) (SEQ ID No. 30)

5'CGCGGATCCTCACACTCTGGTCTCACCCCCACTC3'

so as to amplify a 1589 bp fragment containing the gene encoding the rabies virus G glycoprotein. After purification, the RT-PCR product was digested with PstI and BamHI to give a 1578 bp PstI-BamHI fragment. This fragment was ligated with the vector pVR1012 (Example 6), previously digested with PstI and BamHI, to give the plasmid pAB041 (6437 bp) (Figure No. 16).

Example 20: Production and Purification of the Plasmids

For the preparation of the plasmids intended for the vaccination of animals, any technique may be used that allows for obtaining a suspension of purified plasmids predominantly in the supercoiled form. These techniques are well known to one skilled in the art. There may be mentioned in particular the alkaline lysis technique followed by two successive ultracentrifugations on a caesium chloride gradient in the presence of ethidium bromide as described in J. Sambrook et al. (Molecular Cloning: A Laboratory Manual, 2nd edition, Cold

Spring Harbor Laboratory, Cold Spring Harbor, N.Y., 1989). Reference may also be made to Patent Applications PCT WO 95/21250 and PCT WO 96/02658 which describe methods for producing, on an industrial scale, plasmids which can be used for vaccination. For the purposes of the manufacture of vaccines (see Example 17), the purified plasmids are resuspended so as to obtain solutions at a high concentration (>2 mg/ml) which are compatible with storage. To do so, the plasmids are resuspended either in ultrapure water or in TE buffer (10 mM Tris-HCl; 1 mM EDTA, pH 8.0).

Example 21: Manufacture of the Associated Vaccines

The various plasmids necessary for the manufacture of an associated vaccine are mixed using their concentrated solutions (Example 16). The mixtures are prepared such that the final concentration of each plasmid corresponds to the effective dose of each plasmid. The solutions which can be used to adjust the final concentration of the vaccine may be either a 0.9% NaCl solution, or PBS buffer.

Specific formulations such as liposomes, cationic lipids, may also be used for the manufacture of the vaccines.

Example 22: Vaccination of Cats

The cats are vaccinated with doses of 10 μ g, 50 μ g or 250 μ g per plasmid.

The injections are performed with a needle using the intramuscular route. In that case, the vaccinal doses are administered in a volume of 1 ml.

The injections can also be performed with a needle using the intradermal route. In that case, the vaccinal doses are administered in a total volume of 1 ml administered in 10 injections of 0.1 ml or at 20 injections of 0.05 ml. The intradermal administrations are performed after shaving the skin (thoracic flank in general) or at the level of a relatively glabrous anatomical region, for example the inner surface of the thigh.

A liquid jet injection apparatus (with no needle) can also be used for the intradermal injections.

CLAIMS

1. Vaccine formula intended for cats, comprising at least three polynucleotide vaccine valencies each comprising an integrating plasmid, so as to express in vivo in the host cells, a gene of a feline pathogen valency, these valencies being selected from the group consisting of feline leukaemia virus, panleukopenia virus, infectious peritonitis virus, coryza virus, calicivirovirus, feline immunodeficiency virus and possibly rabies virus, the plasmids comprising, for each valency, one or more of the genes selected from the group consisting of env and gag for the feline leukaemia, VP2 for the panleukopenia, modified S and M for the infectious peritonitis, gB and gD for the coryza, capsid for the calicivirovirus, env and gag/pro for the feline immunodeficiency and G for the rabies.

2. Vaccine formula according to claim 1, characterized in that it comprises the panleukopenia, coryza and calicivirovirus valencies.

3. Vaccine formula according to claim 1 or 2, characterized in that it comprises the coryza virus gB and gD genes, in the same plasmid or in different plasmids, or only one of these genes.

4. Formula according to claim 1, characterized in that it comprises the feline leukaemia virus env and gag genes, in the same plasmid or indifferent plasmids, or the env gene alone.

5. Vaccine formula according to claim 1, characterized in that it comprises the two env and gag/pro genes in different plasmids or in the same plasmid, or only one of these genes.

6. Vaccine formula according to claim 1 or 2, characterized in that it comprises the M gene or the modified S gene in a plasmid or all the genes encoding M or modified S in the same plasmid or in different plasmids.

7. Vaccine formula according to any one of claims 1 to 6, characterized in that it comprises from 10 ng to 1 mg, preferably from 100 ng to 500 µg, still more preferably from 1 µg to 250 µg of each plasmid.

8. Use of one or more plasmids as described in any one of claims 1 to 7, for the manufacture of a vaccine intended to vaccinate cats primo-vaccinated by means of a first vaccine selected from the group consisting of a live whole vaccine, an inactivated whole vaccine, a subunit vaccine, a recombinant vaccine, the first vaccine having the antigen(s) encoded by the plasmid(s) or antigen(s) providing cross-protection.

9. Vaccination kit for cats assembling together a vaccine formula according to any one of claims 1 to 8 and a vaccine for cats selected from the group consisting of a live whole vaccine, an inactivated whole vaccine, a subunit vaccine, a recombinant vaccine, the first vaccine having the antigen encoded by the polynucleotide vaccine or an antigen providing cross-protection, for administration of the latter in primo-vaccination and as a booster with the vaccine formula.

10. Vaccine formula according to any one of claims 1 to 8, accompanied by a notice indicating that this formula can be used as booster for a first vaccine for cats, selected from the group consisting of a live whole vaccine, an inactivated whole vaccine, a subunit vaccine, a recombinant vaccine, this first vaccine having the antigen encoded by the polynucleotide vaccine or an antigen providing cross-protection.

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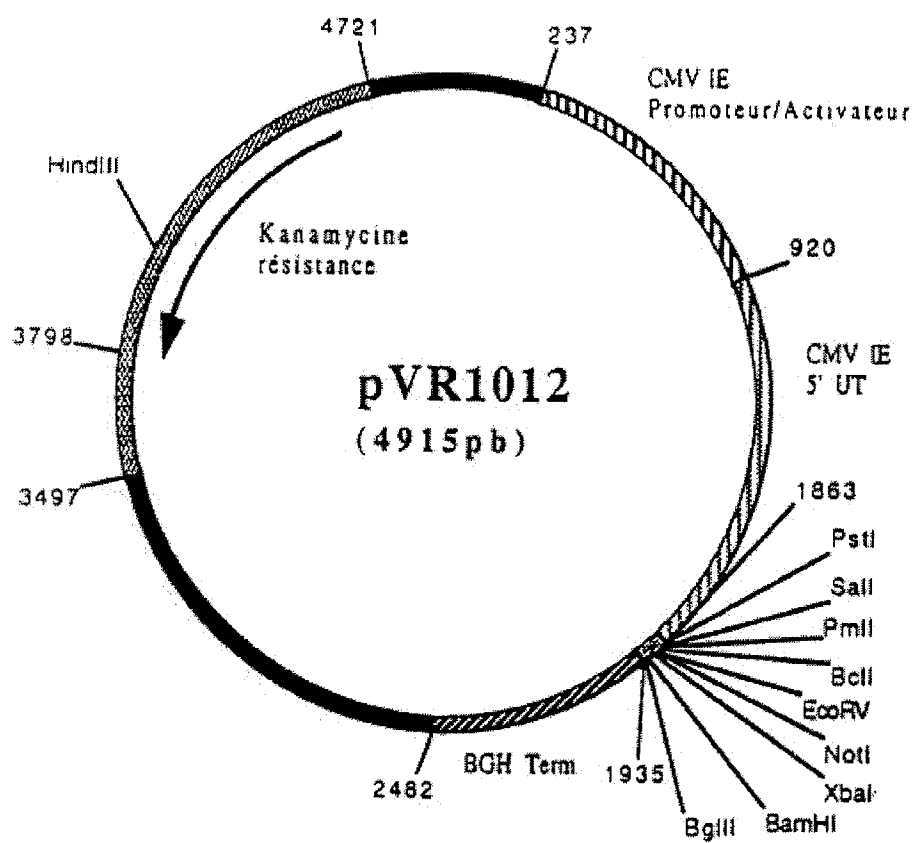


Figure No. 1

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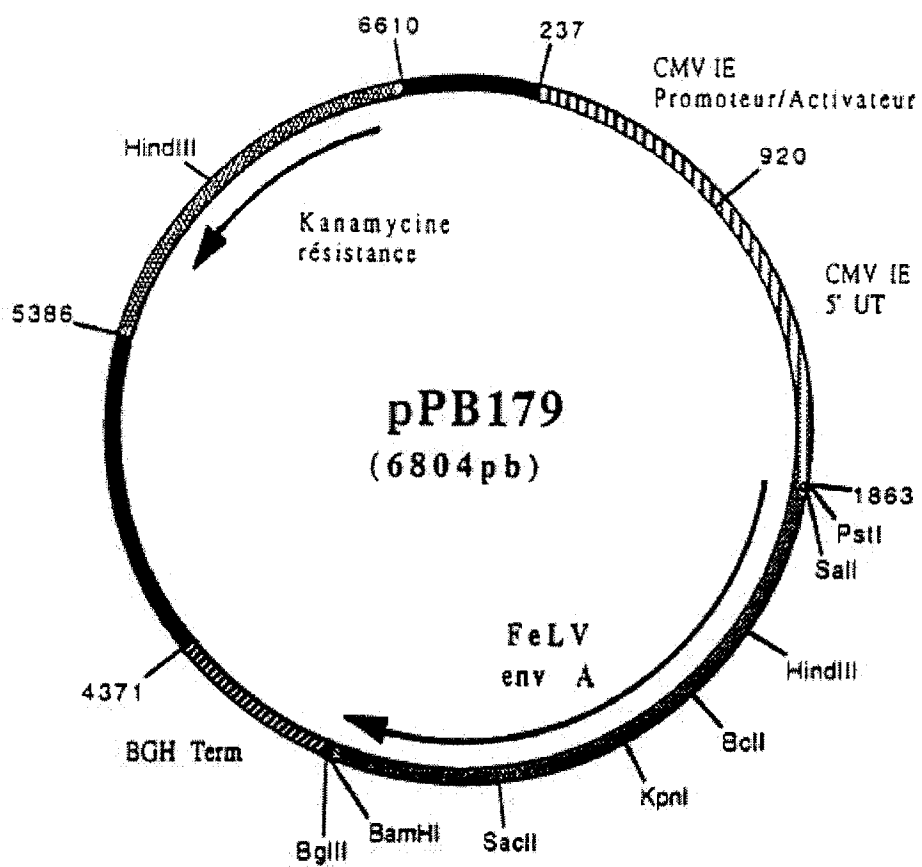


Figure No. 2

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1 ATGGAAGGTCCAACGCACCCAAAACCTCTAAAGATAAGACTTTCTCSTGGGACCTAATGATT
1 MetGluGlyProThrHisProLysProSerLysAspLysThrPheSerTrpAspLeuMetIle

64 CTGCTGGGGCTCTTACTAAGACTGGACGTGGGAATGGCCAATCCTAGTCCGCACCAAATATAT
22 LeuValGlyValLeuLeuArgLeuAspValGlyMetAlaAsnProSerProHisGlnIleTyr

127 AATGTAACCTGGACAATAACCAACCTTGTAAGTGAACAAAGGCTAATGCCACCTCCATGTTG
43 AsnValThrTrpThrIleThrAsnLeuValThrGlyThrLysAlaAsnAlaThrSerMetLeu

190 GGAACCTGACAGACGCTTCCCTACCATGTATTTGACTTATGTGATATAATAGGAAATACA
64 GlyThrLeuThrAspAlaPheProThrMetTyrPheAspLeuCysAspIleIleGlyAsnThr

253 TGGAACCCCTTCAGATCAAGAACCATTCCCAGGGTATGGATGTGATCAGCCTATGAGGAGGTGG
85 TrpAsnProSerAspGlnGluProPheProGlyTyrGlyCysAspGlnProMetArgArgTrp

316 CGACAGAGAAACACACCCCTTTTATGTCTGTCCAGGACATGCCAACCGGAAGCAATGTGGGGGG
106 ArgGlnArgAsnThrProPheTyrValCysProGlyHisAlaAsnArgLysGlnCysGlyGly

379 CCACAGGATGGGTTCGCGCTGTATGGGGTTGCGAGACCACCGGGGAAACCTATTGGAGACCC
127 ProGlnAspGlyPheCysAlaValTrpGlyCysGluThrThrGlyGluThrTyrTrpArgPro

442 ACCTCCTCATGGGACTACATCACAGTAAAAAAGGGGTTACTCAGGGAATATATCAATGTAGT
148 ThrSerSerTrpAspTyrIleThrValLysLysGlyValThrGlnGlyIleTyrGlnCysSer

505 GGAGGTGGTTGGTGTGGGCCCTGTTACGATAAAGCTGTTCACTCCTCGACAACGGGAGCTAGT
169 GlyGlyGlyTrpCysGlyProCysTyrAspLysAlaValHisSerSerThrThrGlyAlaSer

568 GAAGGGGGCCGGTGCACCCCTTGATCTTGCAATTTACCCAAAAGGGAAGACAAACATCTTGG
190 GluGlyGlyArgCysAsnProLeuIleLeuGlnPheThrGlnLysGlyArgGlnThrSerTrp

631 GATGGACCTAAGTCATGGGGCTACGACTATACCGTTCCAGGATATGACCCATAGCCCTGTTTC
211 AspGlyProLysSerTrpGlyLeuArgLeuTyrArgSerGlyTyrAspProIleAlaLeuPhe

694 TCGGTATCCCGCAAGTAATGACCATTACGCCGCTCAGGCCATGGGACCAAATCTAGTCCTG
232 SerValSerArgGlnValMetThrIleThrProProGlnAlaMetGlyProAsnLeuValLeu

757 CCTGATCAAAAACCCCATCCAGGCAATCTCAAATAGAGTCCCGAGTAACACCTCACCATTCC
253 ProAspGlnLysProProSerArgGlnSerGlnIleGluSerArgValThrProHisHisSer

820 CAAGGCAACGGAGGCACCCAGGTGTAACCTTTGTTAATGCCTCCATTGCCCCCTCTACGTACC
274 GlnGlyAsnGlyGlyThrProGlyValThrLeuValAsnAlaSerIleAlaProLeuArgThr

883 CCTGTACCCCCGCAAGTCCCAACGTATAGGGACCGGAAATAGGTTAATAAATTTAGTGCAA
295 ProValThrProAlaSerProLysArgIleGlyThrGlyAsnArgLeuIleAsnLeuValGln

946 GGGACATACCTAGCCTTAAATGCCACCGACCCCAACAAAATAAGACTGTTGGCTCTGCCTG
316 GlyThrTyrLeuAlaLeuAsnAlaThrAspProAsnLysThrLysAspCysTrpLeuCysLeu

1009 GTTCTCGACCACCTTATTACGAAGGGATTGCAATCTTAGGTAACACAGCAACCAACAAAC
337 ValSerArgProProTyrTyrGluGlyIleAlaIleLeuGlyAsnTyrSerAsnGlnThrAsn

1072 CCTCCCCATCTGCCTATCTACTCCGCAACATAAGCTAAGCTATATCTGAGGTGTCAGGGCAA
358 ProSerProSerCysLeuSerThrProGlnHisLysLeuThrIleSerGluValSerGlyGln

1135 GGACTGTGCATAGGGACTGTTCTAAGACCCACCAGGCTTTGTGCAATAAGACACAACAGGGA
379 GlyLeuCysIleGlyThrValProLysThrHisGlnAlaLeuCysAsnLysThrGlnGlnGly

1198 CATACAGGGGCTCACTATCTAGCCGCCCCCAATGGCACCTATTGGGCTGTAACTGGACTC
400 HisThrGlyAlaHisTyrLeuAlaAlaProAsnGlyThrTyrTrpAlaCysAsnThrGlyLeu

Figure No. 3

1261 ACCCCATGCATTTCCATGGCAGTGCCTCAATTGGACCTCTGATTTTTGTGTCTTAATCGAATTA
421> ThrProCysIleSerMetAlaValLeuAsnTrpThrSerAspPheCysValLeuIleGluLeu

1324 TGGCCCAGAGTGACCTACCATCAACCCGAATACATTTACACACATTTCCACAAAGCTGTCAGG
442> TrpProArgValThrTyrHisGlnProGluTyrIleTyrThrHisPheAspLysAlaValArg

1387 TTCCGAAGAGAACCAATATCACTAACC GTT GCCCTTATAATGGGAGGACTCACTGTAGGGGGC
463> PheArgArgGluProIleSerLeuThrValAlaLeuIleMetGlyGlyLeuThrValGlyGly

1450 ATAGCCGCGGGGGTCCGAACAGGGACTAAAGCCCTCCTTGAAACAGCCCAGTTCAGACAATA
484> IleAlaAlaGlyValGlyThrGlyThrLysAlaLeuLeuGluThrAlaGlnPheArgGlnLeu

1513 CAAATGGCTATGCACGCAGACATCCAGGCCCTAGAAGAGTCAATTAGTGCCTTAGAAAAATCC
505> GlnMetAlaMetHisAlaAspIleGlnAlaLeuGluGluSerIleSerAlaLeuGluLysSer

1576 CTGACCTCCCTCTCCGAGGTAGTCTTACAAAATAGACGGGGCCTAGATATTCTGTTCTTACAA
526> LeuThrSerLeuSerGluValValLeuGlnAsnArgArgGlyLeuAspIleLeuPheLeuGln

1639 AAGGGAGGGCTCTGTGCCGCCCTTAAAGGAAGAATGCTGCTTCTATGCAGATCACACCGGACTC
547> LysGlyGlyLeuCysAlaAlaLeuLysGluGluCysCysPheTyrAlaAspHisThrGlyLeu

1702 GTCAGAGACAATATGGCTAAATTAAGAGAAAGACTGAAACAGCGACAACAACCTGTTTGACTCC
568> ValArgAspAsnMetAlaLysLeuArgGluArgLeuLysGlnArgGlnGlnLeuPheAspSer

1765 CAACAGGGATGGTTTGAAGGATGGTTCAACAAGTCCCCCTGGTTTACAACCCTAATTTCTCTCC
589> GlnGlnGlyTrpPheGluGlyTrpPheAsnLysSerProTrpPheThrThrLeuIleSerSer

1828 ATTATAGGCCCCCTTACTAATCCTACTCCTAATTCTCCTCTTCGGCCCCATGCATCCTTAACCGA
610> IleIleGlyProLeuLeuIleLeuLeuLeuIleLeuLeuPheGlyProCysIleLeuAsnArg

1891 TTAGTGCAATTCGTAAAAGACAGAATATCTGTGGTACAAGCCTTAATTTTAACCCAACAGTAC
631> LeuValGlnPheValLysAspArgIleSerValValGlnAlaLeuIleLeuThrGlnGlnTyr

1954 CAACAGATACAGCAATATGATCCGGACCGACCATGA
652> GlnGlnIleGlnGlnTyrAspProAspArgPro...

Figure No. 3 (last part)

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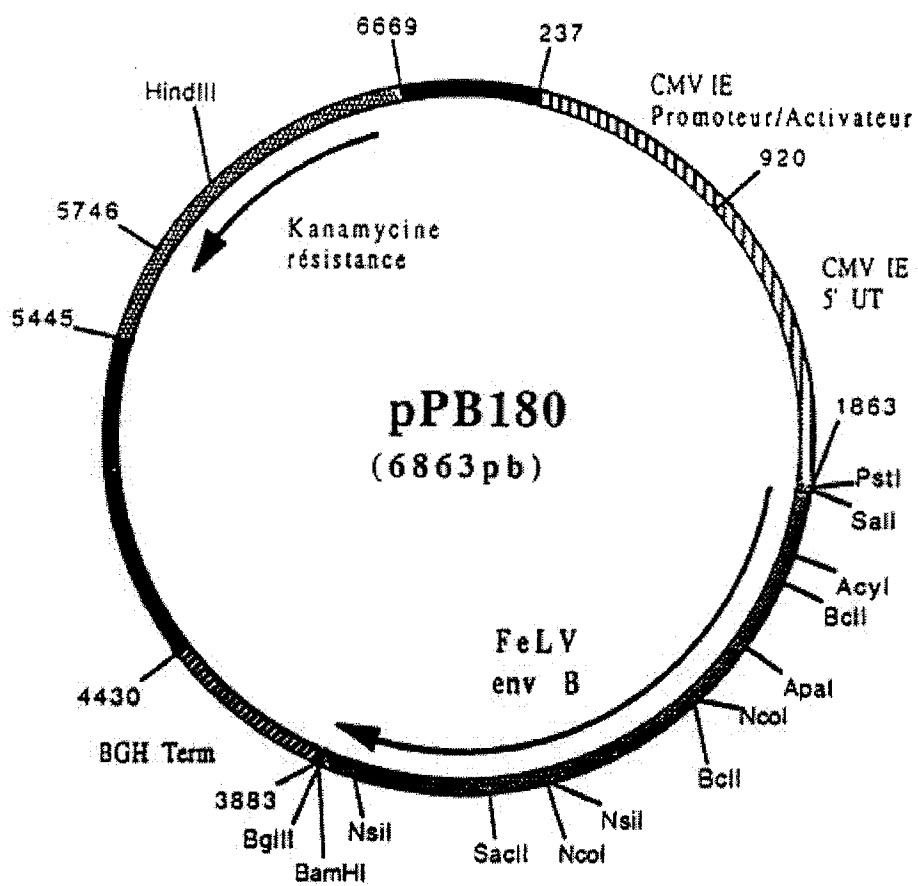


Figure No. 4

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1 ATGCTGGAGCCTCTAGTGGGACAGCCATTGGGGCTCATCTGTTTGGCGTCTCACCTGAATAC
1 Met Ser Gly Ala Ser Ser Gly Thr Ala Ile Gly Ala His Leu Phe Gly Val Ser Pro Glu Tyr

64 AGGTGTTGATCGGAGACGAGGGAGCCGACCCCTCAAGGTCTCTTTCTGAGGTTTCATTTTCG
22 Arg Val Leu Ile Gly Asp Glu Gly Ala Gly Pro Ser Arg Ser Leu Ser Glu Val Ser Phe Ser

127 GTTTGGTACCAAAGACGCGCGGCACGTCTTGTCTATTTTGTCTGTTGCGTCTTTTCTTGTC
43 Val Trp Tyr Gln Arg Arg Ala Ala Arg Leu Val Ile Phe Cys Leu Val Ala Ser Phe Leu Val

190 CCTTGTCTAACCTTTTAAATTGCAGAAACCGTCATGGGCCAACTATAACTACCCCTTAAGC
64 Pro Cys Leu Thr Phe Leu Ile Ala Glu Thr Val Met Gly Gln Thr Ile Thr Thr Pro Leu Ser

253 CTCACCCCTTGATCACTGGTCTGAAGTCCGGGCACGAGCCCATATCAAGGTGTGAGGTCCGG
85 Leu Thr Leu Asp His Trp Ser Glu Val Arg Ala Arg Ala His Asn Gln Gly Val Glu Val Arg

316 AAAAAGAAATGGATTACCTTATGTGAGGCCGAATGGGTGATGATGAATGTGGGCTGGCCCCGA
106 Lys Lys Lys Trp Ile Thr Leu Cys Glu Ala Glu Trp Val Met Met Asn Val Gly Trp Pro Arg

379 GAAGGAACCTTTTCTCTTGATAGCATTTCACAGGTTGAAAAGAAGATCTTCGCCCCGGGACCA
127 Glu Gly Thr Phe Ser Leu Asp Ser Ile Ser Gln Val Glu Lys Lys Ile Phe Ala Pro Gly Pro

442 TATGGACACCCCGACCAAGTTCCTTACATTACTACATGGAGATCCTTAGCCACAGACCCCCCT
148 Tyr Gly His Pro Asp Gln Val Pro Tyr Ile Thr Thr Trp Arg Ser Leu Ala Thr Asp Pro Pro

505 TCGTGGGTTCGTCCGTTCCCTACCCCTCCCAAACCTCCACACCCCTCCCTCAACCTCTTTTCG
169 Ser Trp Val Arg Pro Phe Leu Pro Pro Pro Lys Pro Pro Thr Pro Leu Pro Gln Pro Leu Ser

568 CCGCAGCCCTCCGCCCCCTTACCTCTTCCCTCTACCCCGTTCTCCCCAAGCCAGACCCCCC
190 Pro Gln Pro Ser Ala Pro Leu Thr Ser Ser Leu Tyr Pro Val Leu Pro Lys Pro Asp Pro Pro

631 AAACCGCTGTGTTACCGCCTGATCCTTCTTCCCTTTAATTGATCTCTTAACAGAAGAGCCA
211 Lys Pro Pro Val Leu Pro Pro Asp Pro Ser Ser Pro Leu Ile Asp Leu Leu Thr Glu Glu Pro

694 CCTCCCTATCCGGGGGGTCACGGGCCACCGCCATCAGGTCCTAGGACCCCAACCGCTTCCCCG
232 Pro Pro Tyr Pro Gly Gly His Gly Pro Pro Pro Ser Gly Pro Arg Thr Pro Thr Ala Ser Pro

757 ATTGCAAGCCGGCTAAGGGAACGACGAGAAAACCTGCTGAAGAATCGCAAGCCCTCCCTTGG
253 Ile Ala Ser Arg Leu Arg Glu Arg Arg Glu Asn Pro Ala Glu Glu Ser Gln Ala Leu Pro Leu

820 AGGGAAGGCCCCAACAACCGACCCAGTATTGGCCATTCTCAGCTTCAGACTTGTATAACTGG
274 Arg Glu Gly Pro Asn Asn Arg Pro Gln Tyr Trp Pro Phe Ser Ala Ser Asp Leu Tyr Asn Trp

883 AAGTCGCATAACCCCTTTCTCCCAAGATCCAGTGGCCCTAACTAACCTAATTGAGTCCATT
295 Lys Ser His Asn Pro Pro Phe Ser Gln Asp Pro Val Ala Leu Thr Asn Leu Ile Glu Ser Ile

946 TTAGTGACGCATCAACCAACCTGGGACGACTGCCAGCAGCTCTTGCAGGCACTCCTGACAGGC
316 Leu Val Thr His Gln Pro Thr Trp Asp Asp Cys Gln Gln Leu Leu Gln Ala Leu Leu Thr Gly

1009 GAAGAAAGGCAAAGGGTCTTCTTGGAGCCCGAAAGCAGGTTCCAGGCGAGGACGGACGGCCA
337 Glu Glu Arg Gln Arg Val Leu Leu Glu Ala Arg Lys Gln Val Pro Gly Glu Asp Gly Arg Pro

1072 ACCCAACTACCAATGTCTATTGACGAGACTTTCCCTTGACCCGTCCTCAACTGGGATTTTGCT
358 Thr Gln Leu Pro Asn Val Ile Asp Glu Thr Phe Pro Leu Thr Arg Pro Asn Trp Asp Phe Ala

1135 ACGCCGGCAGGTAGGGAGCACCTACGCCTTTATCGCCAGTTGCTATTAGCGGCTCTCCGCGG
379 Thr Pro Ala Gly Arg Glu His Leu Arg Leu Tyr Arg Gln Leu Leu Leu Ala Gly Leu Arg Gly

Figure No. 5

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1198 GCTGCAAGACGCCCCACTAATTTGGCACAGGTAAAGCAGGTTGTACAAGGGAAGAGGAAACG
 400▶ AlaAlaArgArgProThrAsnLeuAlaGlnValLysGlnValValGlnGlyLysGluGluThr
 1261 CCAGCAGCATTTTTAGAAAGATTAAAAGAGGCTTATAGAATGTACACTCCCTATGACCCTGAG
 421▶ ProAlaAlaPheLeuGluArgLeuLysGluAlaTyrArgMetTyrThrProTyrAspProGlu
 1324 CACCCAGGGCAAGCGGCTAGTGTATCCTATCCTTTATATACCACTCTAGCCAGATATAAGA
 442▶ AspProGlyGlnAlaAlaSerValIleLeuSerPheIleTyrGlnSerSerProAspIleArg
 1387 AATAAGTTACAAAGGCTAGAAGGCCTACAAGGGTTCACCCTATCTGATCTGCTAAAAGAGGCA
 463▶ AsnLysLeuGlnArgLeuGluGlyLeuGlnGlyPheThrLeuSerAspLeuLeuLysGluAla
 1450 GAAAAGATATACAACAAAAGGGAGACCCAGAGGAAAGGGAAGAAAGATTATGGCAGCGACAG
 484▶ GluLysIleTyrAsnLysArgGluThrProGluGluArgGluGluArgLeuTrpGlnArgGln
 1513 GAAGAAAGAGATAAAAAGCGCCACAAGGAGATGACTAAAGTTCTGGCCACAGTAGTTGCTCAG
 505▶ GluGluArgAspLysLysArgHisLysGluMetThrLysValLeuAlaThrValValAlaGln
 1576 AATAGAGATAAGGATAGAGAAGAAAGTAACTGGGGGATCAAAGGAAAATACCTCTGGGGAAA
 526▶ AsnArgAspLysAspArgGluGluSerLysLeuGlyAspGlnArgLysIleProLeuGlyLys
 1639 GACCACTGTGCCTATTGCAAGGAAAAGGGGCATTGGGTTCCGCGATTGCCCCAAACGACCCAGG
 547▶ AspGlnCysAlaTyrCysLysGluLysGlyHisTrpValArgAspCysProLysArgProArg
 1702 AAGAAACCCGCCAACTCCACTCTCCTCAACTTAGGAGATTAGGAGAGTCAGGGCCAGGACCCC
 568▶ LysLysProAlaAsnSerThrLeuLeuAsnLeuGlyAsp...
 1▶ GluIleArgArgValArgAlaArgThrPr
 1765 CCCCCCTGAGCCCAGGATAACCTTAAAAATAGGGGGGCAACCGGTGACTTTTCTGGTGGAC
 10▶ oProProGluProArgIleThrLeuLysIleGlyGlyGlnProValThrPheLeuValAspTh
 1828 GGGAGCCCAGCACTCAGTACTGACTCGACCAGATGGACCTCTCAGTGACCGCACAGCCCTGGT
 31▶ rGlyAlaGlnHisSerValLeuThrArgProAspGlyProLeuSerAspArgThrAlaLeuVa
 1891 GCAAGGAGCCACGGGAAGCAAAAACCTACCGGTGGACCACCGACAGGAGGGTACAACCTGGCAAC
 52▶ lGlnGlyAlaThrGlySerLysAsnTyrArgTrpThrThrAspArgArgValGlnLeuAlaTh
 1954 CGGTAAGGTGACTCATTCTTTTTTATATGTACCTGAATGTCCCTACCCGTTATTAGGGAGAGA
 73▶ rGlyLysValThrHisSerPheLeuTyrValProGluCysProTyrProLeuLeuGlyArgAs
 2017 CCTATTAACTAACTTAAGGCCCAATCCATTTTACCGGAGAAGGGGCTAATGTTGTTGGGCC
 94▶ pLeuLeuThrLysLeuLysAlaGlnIleHisPheThrGlyGluGlyAlaAsnValValGlyPr
 2080 CAGGGGTTTACCCCTACAAGTCTTACTTTACAATTAGAAGAGGAGTATCGGCTATTTGAGCC
 115▶ oArgGlyLeuProLeuGlnValLeuThrLeuGlnLeuGluGluGluTyrArgLeuPheGluPr
 2143 AGAAAGTACACAAAACAGGAGATGGACACTTGGCTTAAAACTTTCCCAGGCGTGGGCAGA
 136▶ oGluSerThrGlnLysGlnGluMetAspThrTrpLeuLysAsnPheProGlnAlaTrpAlaGl

Figure No. 5 (middle part)

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2206 AACAGGAGGTATGGGAATGGCTCATTGTCAAGCCCCCTTCTCATTCAACTTAAGGCTACTGC
157>uThrGlyGlyMetGlyMetAlaHisCysGlnAlaProValLeuIleGlnLeuLysAlaThrAl
2269 CACTCCAATCTCCATCCGACAGTATCCTATGCCCCATGAAGCGTACCAGGGAATTAAGCCTCA
178>aThrProIleSerIleArgGlnTyrProMetProHisGluAlaTyrGlnGlyIleLysProHi
2332 TATAAGAAGAATGCTAGATCAAGGCATCCTCAAGCCCTGCCAGTCCCCATGGAATACACCCTT
199>sIleArgArgMetLeuAspGlnGlyIleLeuLysProCysGlnSerProTrpAsnThrProLe
2395 ATTACCTGTTAAGAAGCCAGGGACCGAGGATTACAGACCAGTGCAGGACTTAAGAGAAGTAAA
220>uLeuProValLysLysProGlyThrGluAspTyrArgProValGlnAspLeuArgGluValAs
2458 CAAAAGAGTAGAAGACATCCATCCTACTGTGCCAAATCCATATAACCTCCTTAGCACCCCTCCC
241>nLysArgValGluAspIleHisProThrValProAsnProTyrAsnLeuLeuSerThrLeuPr
2521 GCCGTCTCACCCTTGGTACACTGTCCTAGATTTAAAGGACGCTTTTTTCTGCCTGCGACTACA
262>oProSerHisProTrpTyrThrValLeuAspLeuLysAspAlaPhePheCysLeuArgLeuHi
2584 CTCTGAGAGTCAGTTACTTTTTGCATTTGAATGGAGAGATCCAGAAATAGGACTGTCAGGGCA
283>sSerGluSerGlnLeuLeuPheAlaPheGluTrpArgAspProGluIleGlyLeuSerGlyGl
2647 ACTAACCTGGACACGCCTTCCTCAGGGGTCAAGAATAGCCCCACCCTATTTGATGAGGGCCCT
304>nLeuThrTrpThrArgLeuProGlnGlyPheLysAsnSerProThrLeuPheAspGluAlaLe
2710 GCACTCAGACCTGGCCGATTTTCAGGGTAAGGTACCCGGCTCTAGTCCTCCTACAATATGTAGA
325>uHisSerAspLeuAlaAspPheArgValArgTyrProAlaLeuValLeuLeuGlnTyrValAs
2773 TGACCTCTTGCTGGCTGCGGCAACCAGGACTGAATGCCTGGAAGGGACTAAGGCACTCCTTGA
346>pAspLeuLeuLeuAlaAlaAlaThrArgThrGluCysLeuGluGlyThrLysAlaLeuLeuGl
2836 GACTTTGGGCAATAAGGGGTACCGAGCCTCTGGAAAGAAGGCCCAAATTTGCCTGCAAGAAGT
367>uThrLeuGlyAsnLysGlyTyrArgAlaSerGlyLysLysAlaGlnIleCysLeuGlnGluVa
2899 CACATACCTGGGTACTCTTTAAAAGATGGCCAAAGGTGGCTTACCAAAGCTCGGAAAGAAGC
388>lThrTyrLeuGlyTyrSerLeuLysAspGlyGlnArgTrpLeuThrLysAlaArgLysGluAl
2962 CATCCTATCCATCCCTGTGCCTAAAAACCCACGACAAGTGAGAGAGTTCTTGGAACTGCAG
409>aIleLeuSerIleProValProLysAsnProArgGlnValArgGluPheLeuGlyThrAla

Figure No. 5 (Last part)

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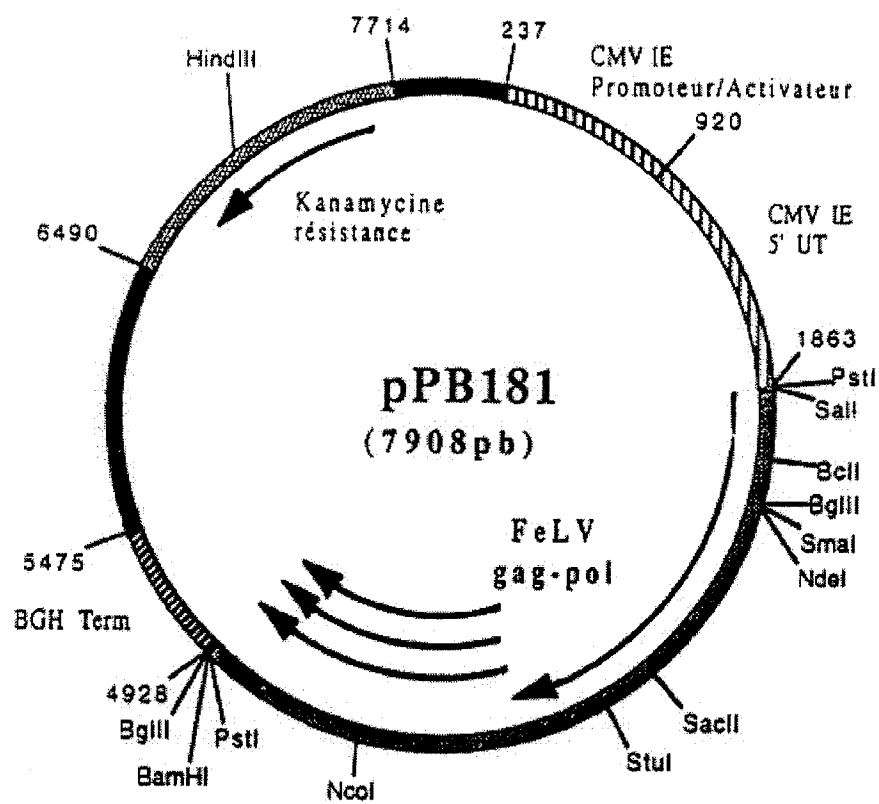


Figure No. 6

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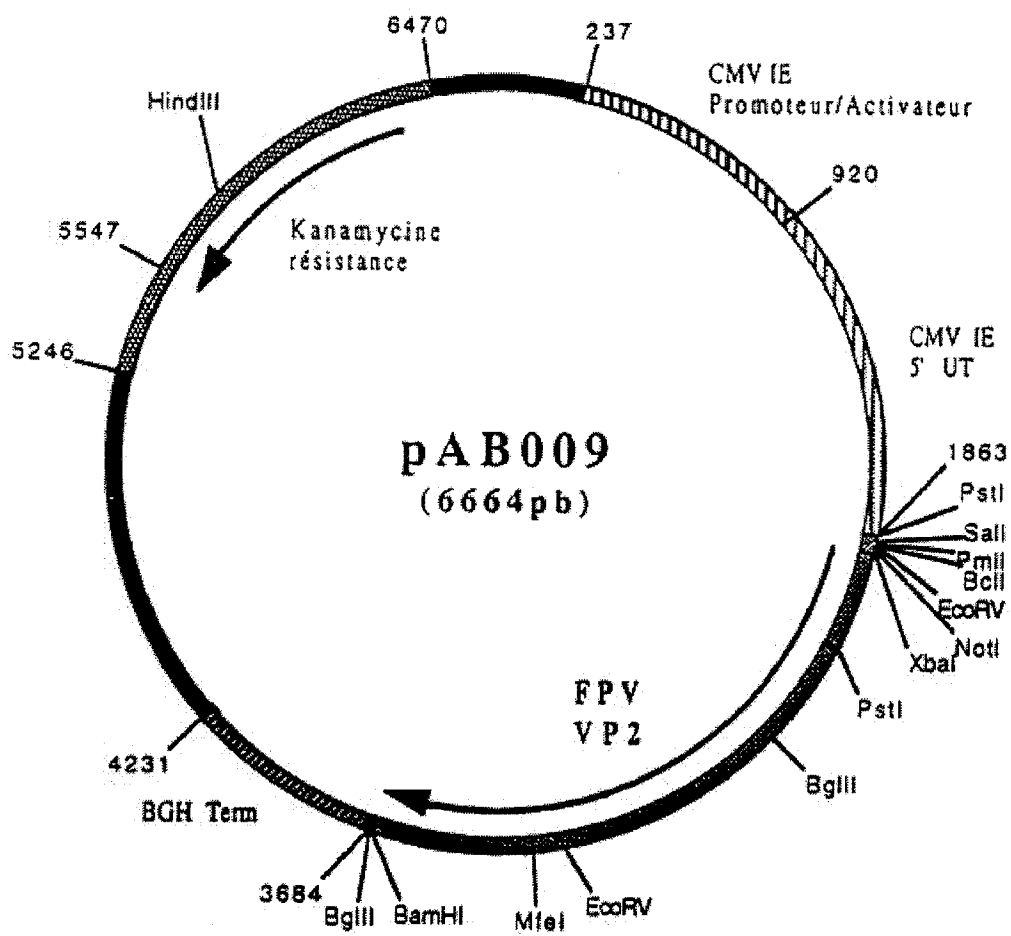


Figure No. 7

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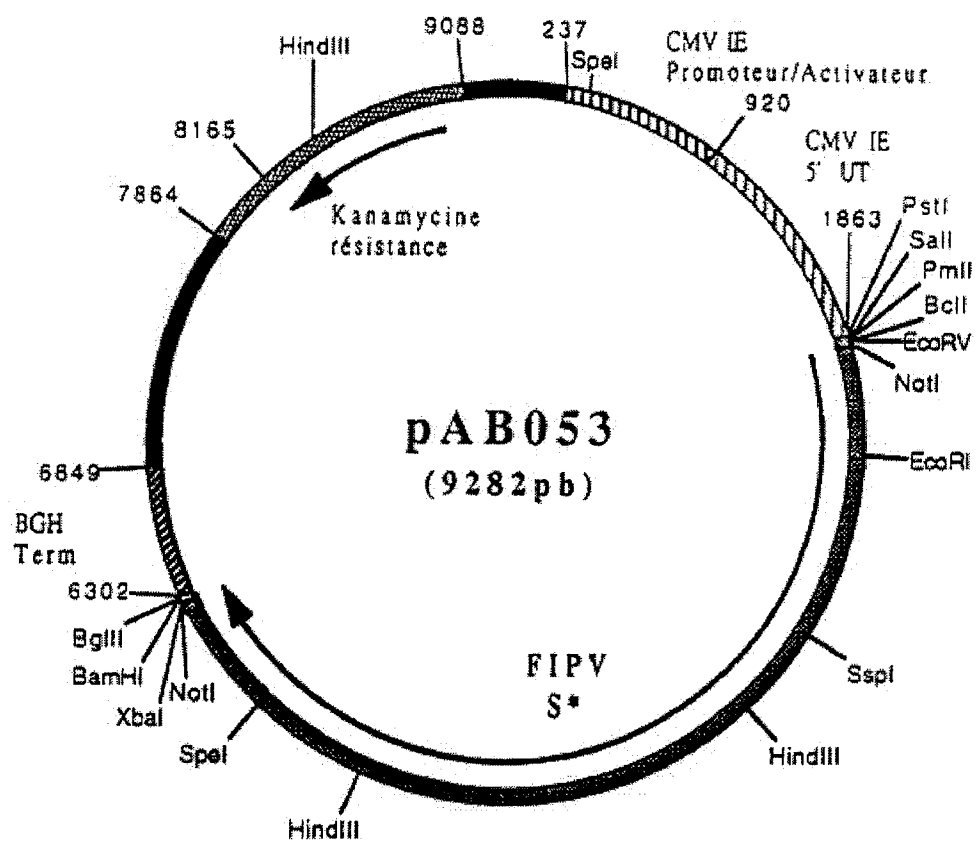


Figure No. 8

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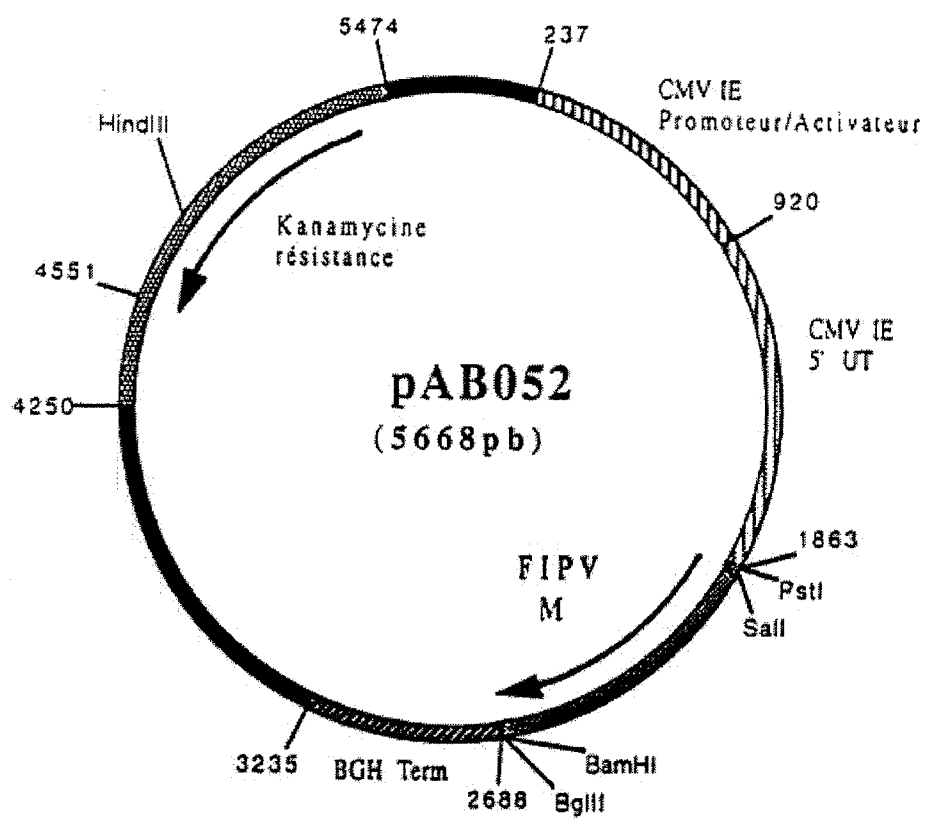


Figure No. 9

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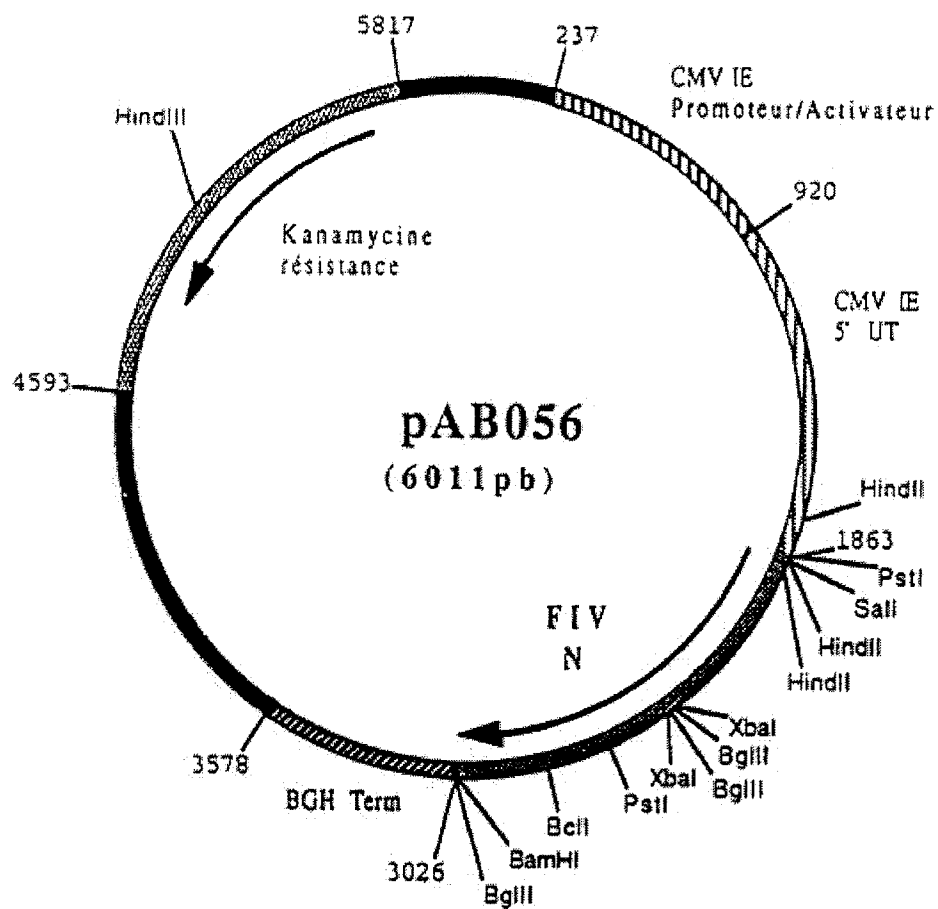


Figure No. 10

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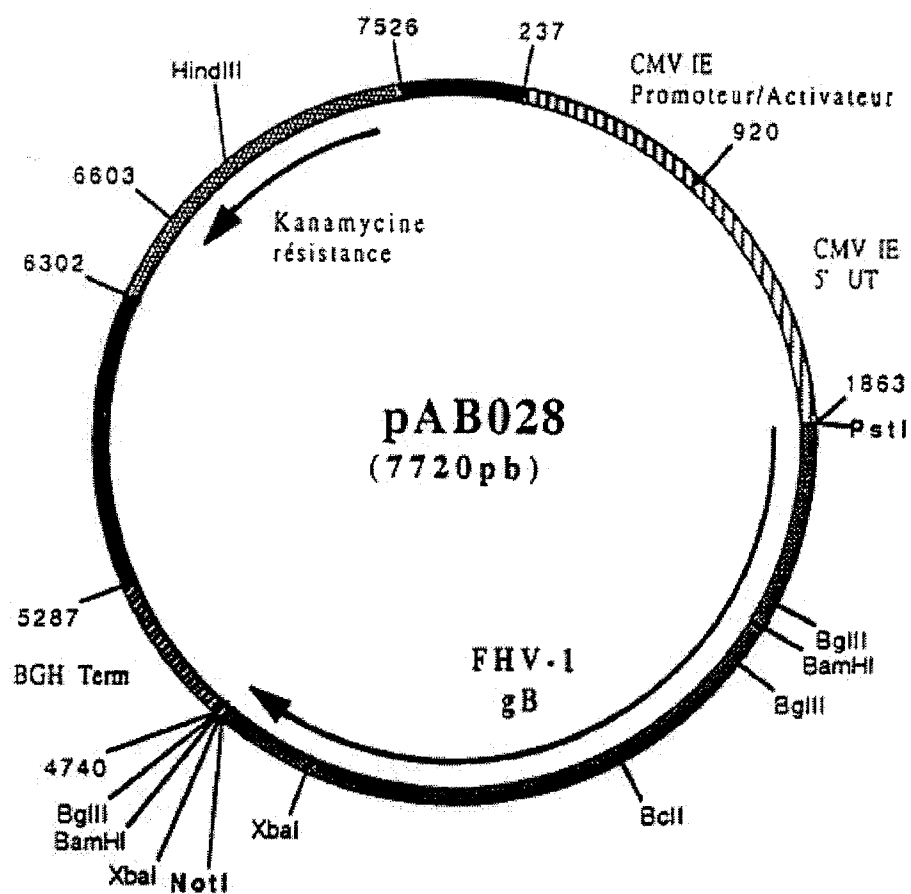


Figure No. 11

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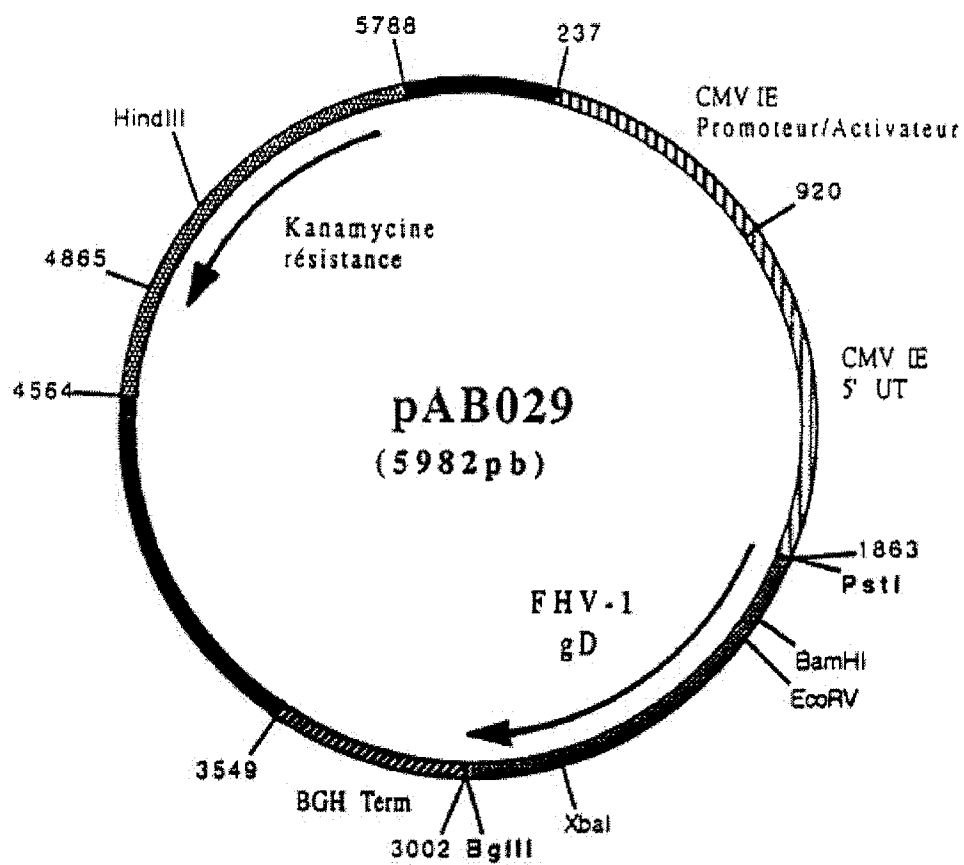


Figure No. 12

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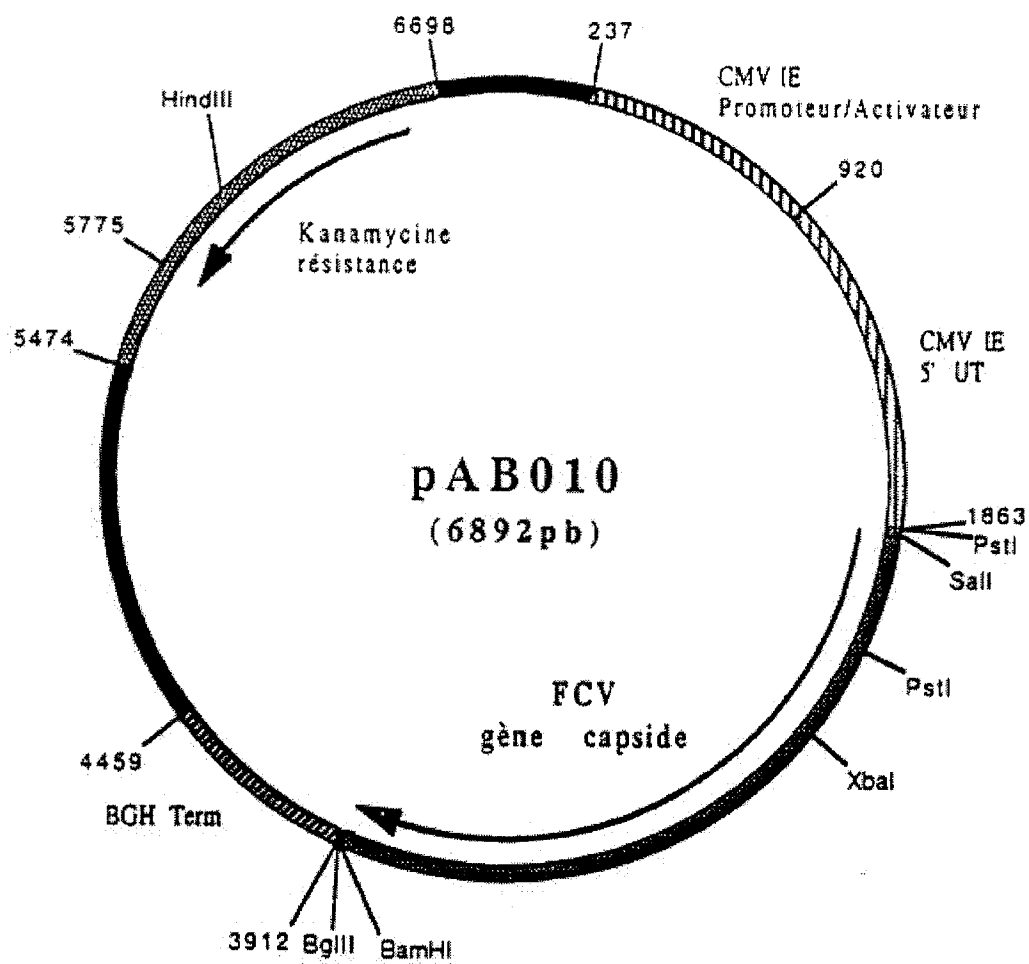


Figure No. 13

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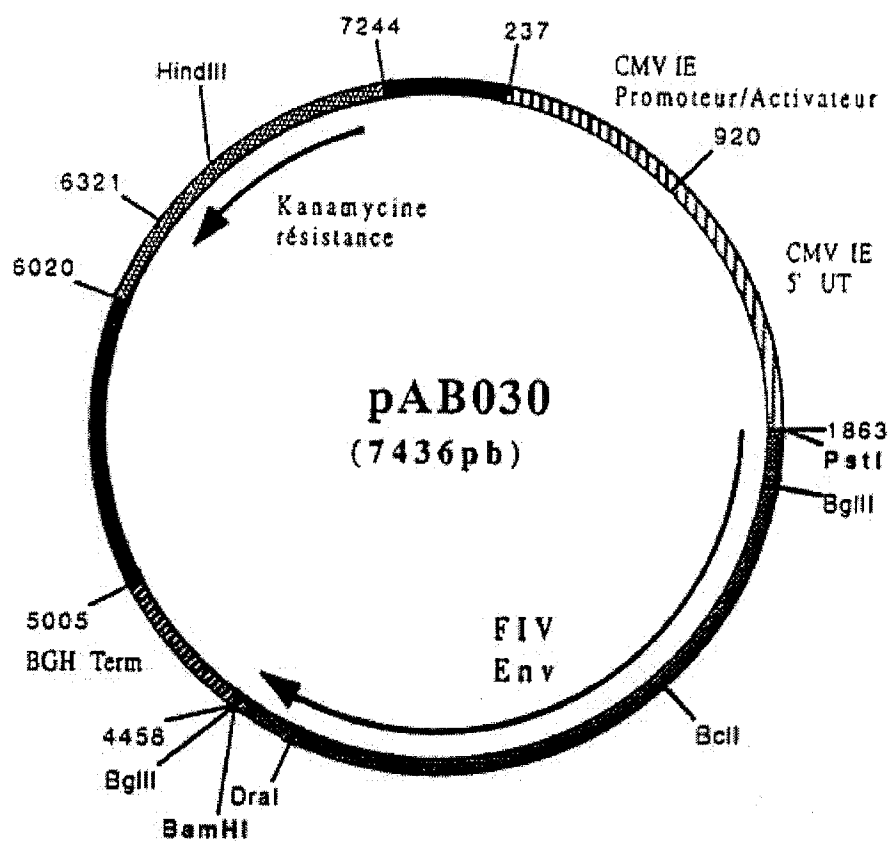


Figure No. 14

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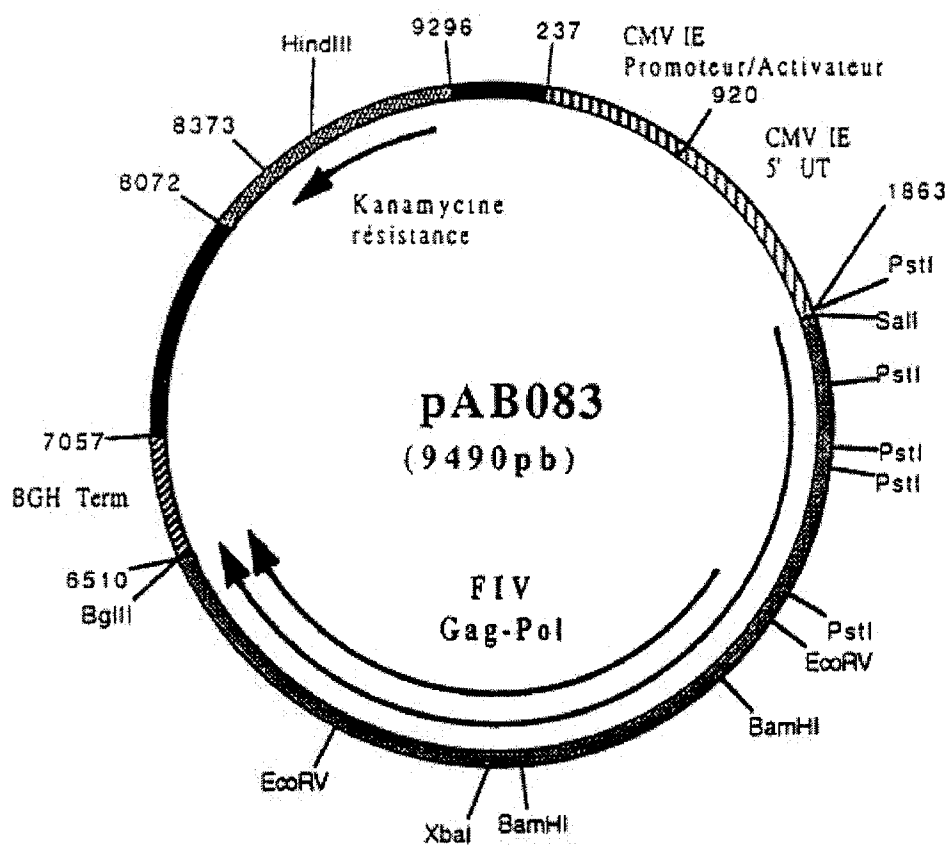


Figure No. 15

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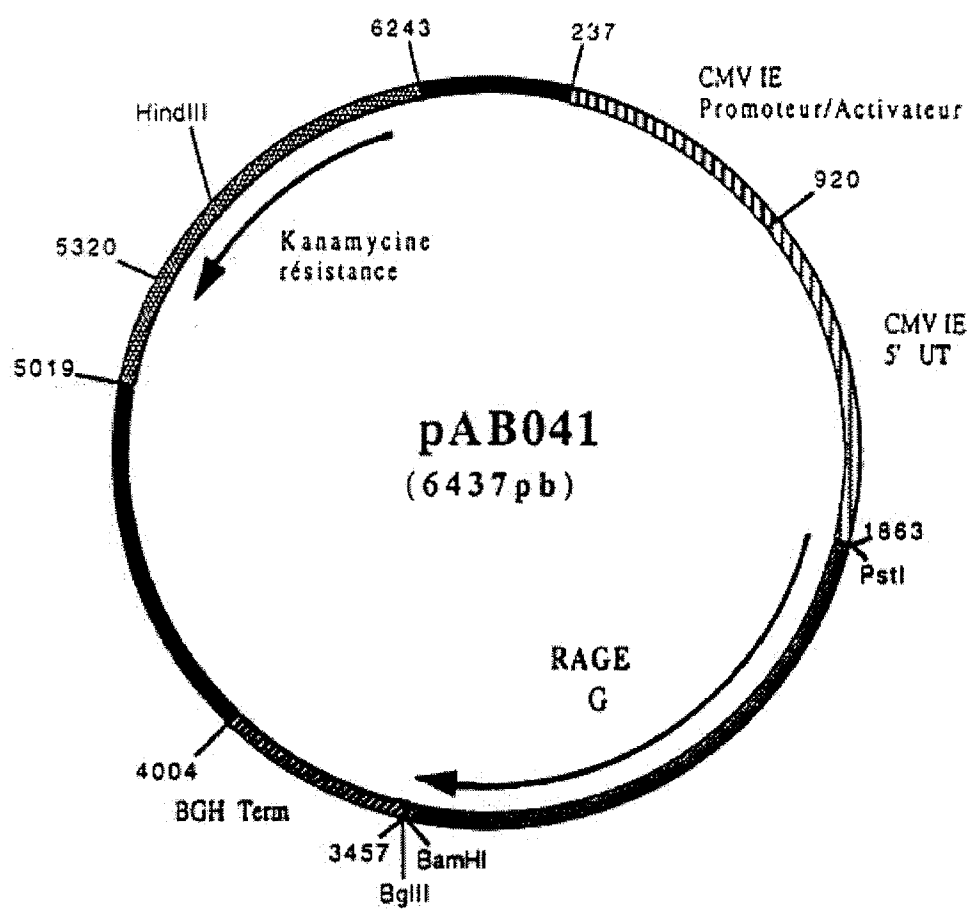


Figure No. 16